

PCT09

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/462,972

DATE: 02/14/2003 \$6

TIME: 11:52:50

Input Set : A:\BB-1095-A Corrected Seq List.txt Output Set: N:\CRF4\02142003\I462972.raw

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61

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Input Set : A:\BB-1095-A Corrected Seq List.txt Output Set: N:\CRF4\02142003\1462972.raw

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	452
65 115 att ctt cac tta gtt ctg cca age Lys Val Asp Pro	
64 Asn Ile Asp Ile Thr Lys IIe III 120 65 115 120 67 aac gac agt tat gtt ctt cac tta gtt ctg cca agt aaa gtt gat cca 68 Asn Asp Ser Tyr Val Leu His Leu Val Leu Pro Ser Lys Val Asp Pro 68 Asn Asp Ser Tyr Val Leu His Leu Val Leu Pro Ser Lys Val Asp Pro 135 130 130 135	480
125 125 ASD DC4 -1	400
67 add gd	
71 ttg gtc aat ctg atg the Tys Val Glu Lys Val F10 MF 160	528
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73 145 agt tta gac cag caa att ada gad Lys Glu Val 11e	
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75 atg det 77 76 Met Ile Gly Gly Leu Asp GIN GIN 170 77 79 gag cta cca atc aaa cat cct gag ctg ttt gaa agt ctt gga att gca 77 79 gag cta cca atc aaa cat cct gag ctg ttt gaa agt ctt gga att gca 79 79 gag cta cca atc aaa cat cct gag ctg ttt gaa agt ctt gga att gca 79 79 gag cta cca atc aaa cat cct gag ctg ttt gaa agt ctt gga att gca 79 79 gag cta cca atc aaa cat cct gag cca cct ggt aca gga aaa aca 780	
79 gag cta coa aco His Pro Glu Leu File 3190	624
81 and get get cet get tat ggg cod Pro Gly Thr Gly Lys Int	
80 Glu Leu Pro Ile Lys His Pro Ile 185 81 180 82 caa cca aag ggt gtc ctg ctc tat ggg cca cct ggt aca ggt aaa aca 83 caa cca aag ggt gtc ctg ctc tat ggg cca cct ggt aca ggt aaa aca 84 Gln Pro Lys Gly Val Leu Leu Tyr Gly Pro Pro Gly Thr Gly Lys Thr 84 Gln Pro Lys Gly Val Leu Leu Tyr Gly Pro Pro Gly Thr Gly Lys Thr 85 205 86 Glu Leu Pro Ile Lys His Pro Ile 185 87 Cat act gac tgt aca ttc atc agg	672
84 GHR 195 85 195 87 ttg ttg gct agg gca gtg gct cat cat act gac tgt aca ttc duby 88 Leu Leu Ala Arg Ala Val Ala His His Thr Asp Cys Thr Phe Ile Arg 88 Leu Leu Ala Arg Ala Val Ala His His Thr Asp Cys Thr Phe Ile Arg 88 Leu Leu Ala Arg Ala Val Ala His His Thr Asp Cys Thr Phe Ile Arg 88 Leu Leu Ala Arg Ala Val Ala His His Thr Asp Cys Thr Phe Ile Arg 88 Leu Leu Ala Arg Ala Val Ala His His Thr Asp Cys Thr Phe Ile Arg 88 Leu Leu Ala Arg Ala Val Ala His His Thr Asp Cys Thr Phe Ile Arg 88 Leu Leu Ala Arg Ala Val Ala His His Thr Asp Cys Thr Phe Ile Arg	
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88 Led led 112 215 216 216 and tac att gga gaa ggt tee as	
88 Leu Leu Ala Arg Ala Val Ala III 220 88 Leu Leu Ala Arg Ala Val Ala III 220 89 210 91 gtg tct ggt tct gag tta gtt cag aaa tac att gga gaa ggt tct aga 91 gtg tct ggt tct gag tta gtt cag aaa tac att gga gaa ggt tct aga 92 Val Ser Gly Ser Glu Leu Val Gln Lys Tyr Ile Gly Glu Gly Ser Arg 92 Val Ser Gly Ser Glu Leu Val Gln Lys Tyr Ile Gly Glu Gly Ser Arg 92 Val Ser Gly Ser Glu Leu Val Gln Lys Tyr Ile Gly Glu Gly Ser Arg 92 Val Ser Gly Ser Glu Leu Val Gln Lys Tyr Ile Gly Glu Gly Ser Arg 93 Val Ser Gly Ser Glu Leu Val Gln Lys Tyr Ile Gly Glu Gly Ser Arg	
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* 97 245 att gac agt att gga tct gct cgg ats Ser	
• 99 atc ttc atg gat gaa atc gab Ser Ile Gly Ser Ala 123 270	864
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100 The Thomas 260 101 260 103 gga agt ggc aac ggt gat agt gag gta cag cgt act atg ctg gdd 103 gga agt ggc aac ggt gat agt gag gta cag cgt act atg ctg gdd 104 Gly Ser Gly Asn Gly Asp Ser Glu Val Gln Arg Thr Met Leu Glu Leu 285 280 280 275 275 275 280 280 275	
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104 Gly Ser Gly Ash Gly and 280	712
105 275 ttg gat gga ttt gaa gct tca aac ary Ile Lys Val Leu	
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115 gga cgg acc s Lys Ile Glu File 22	1056
116 GIV ALY 123 325 and tot aga aga atg aat tta acy og	
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129 370 The cac gtg act cag gag gat ttt gag des Ala Val Ala	
128 Ala Glu Leu Lys Ala Val Gys 375 129 370 131 gag cgg agg gta cac gtg act cag gag gat ttt gag atg gcc gtg gcg 131 gag cgg agg gta cac gtg act cag gag gat ttt gag atg gcc gtg gcg 131 gag cgg agg gta cac gtg act cag gag gat ttt gag atg gcc gtg gcg 132 Glu Arg Arg Val His Val Thr Gln Glu Asp Phe Glu Met Ala Val Ala 395 390 390 390 380 390 400	1248
122 GIII ALU ALU ALU AND	1240
132 Glu Arg Arg Val His Val His 395 133 385 135 aag gtg atg aaa aag gag act gaa aaa aac atg tca ttg cgg aag ttg 135 aag gtg atg aaa aag gag act gaa aaa aac atg tca ttg cgg aag ttg 136 Lys Val Met Lys Lys Glu Thr Glu Lys Asn Met Ser Leu Arg Lys Leu 136 Lys Val Met Lys Lys Glu Thr Glu Lys Asn Met Ser Leu Arg Lys Lys 415	
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156 To The His Giu His of As	
159 His Tyr Tyr Ser Leu Ash 116 40 160 35 40 Ala Gln Arg Ash Asp Leu Ash 160 55 60 60 60 60 60 60 60 60 60 60 60 60 60	
160 Asn Leu Asn Arg Leu Glu Pro	
163 Lys Thr His Asn Leu Asn Arg Bob 60 163 Lys Thr His Asn Leu Asn Arg Bob 60 164 50 55 164 50 75 167 Ser Arg Val Arg Met Leu Arg Glu Glu Leu Gln Leu Leu Gln Glu Pro 75 70 70 111 Lys Val Met Gly Lys Asn Lys Val	
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196 205 200 200 Pro Lys Gly Val Leu Leu Tyl Gly 200 205	វ
200 - 77 7 712 HIS HIS THE 77 7	
200 Leu Leu Ala Arg Ala Val Ala Mis 220 220 215 215 220 220 220 220 220 220 220 220 220 22	3
204 210 Tai Val GIN LVS 11 - 24	U
203 Leu Leu Ala Arg Ala Val Ala III 220 204 210 207 Val Ser Gly Ser Glu Leu Val Gln Lys Tyr Ile Gly Glu Gly Ser Arg 208 225 211 Met Val Arg Glu Leu Phe Val Met Ala Arg Glu His Ala Pro Ser Il 255 245 245 250 267 Jlo Gly Ser Ala Arg Met Glu Se	е
208 225 $pho Val Met Ala Ala = 255$	
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243 Ala Glu Bour 1 375 Asp Phe Glu Met Ala Vai	
243 Ala Glu Leu Lys Ala Val Cys III 380 243 Ala Glu Leu Lys Ala Val Cys III 380 244 370 247 Glu Arg Arg Val His Val Thr Gln Glu Asp Phe Glu Met Ala Val Ala 390 247 Glu Arg Arg Val His Val Thr Gln Glu Asp Phe Glu Met Ala Val Ala 390 390 390 390 390 390 390 390 390 390	
244 San Arg Val His Val Thr Grives 395 247 Glu Arg Arg Val His Val Thr Grives 395 248 385 251 Lys Val Met Lys Lys Glu Thr Glu Lys Asn Met Ser Leu Arg Lys Leu 415	
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271 GIU HIS 120 5 5 and car car gac ctc aac tcc cga yee	
271 Glu His Ile His Asp Leu Glu 10 272 1 274 aac ctc aac cgc ctc gag gcc cag cgc aac gac ctc aac tcc cga gtt 275 Asn Leu Asn Arg Leu Glu Ala Gln Arg Asn Asp Leu Asn Ser Arg Val 275 20 276 277 278 278 279 279 279 279 279 279 279 279 279 279	4.12
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291 The Third 85 292 80 292 Roman Street S	
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312 160 314 cgt gcg gtt gct cat cac act gac tgc acc ttc atc agg gtg tct ggt 315 Arg Ala Val Ala His His Thr Asp Cys Thr Phe Ile Arg Val Ser Gly 316 318 tct gag ttg gtt cag aag tat att ggt gag ggc tcc cgg atg gtt agg 319 Ser Glu Leu Val Gln Lys Tyr Ile Gly Glu Gly Ser Arg Met Val Arg 319 Ser Glu Leu Val ggc agg gaa cat gca cca tcc att ata ttt atg 312 gaa ctc ttt gtt atg gcc agg gaa cat gca cca tcc att ata ttt atg 312 agg act ctt gtt atg gcc agg gaa cat gca cca tcc att ata ttt atg 312 aga ctc ttt gtt atg gcc agg gaa cat gca cca tcc att ata ttt atg 312 aga ctc ttt gtt atg gcc agg gaa cat gca cca tcc att ata ttt atg 312 aga ctc ttt gtt atg gcc agg gaa cat gca cca tcc att ata ttt atg 312 aga ctc ttt gtt atg gcc agg gaa cat gca cca tcc att ata ttt atg 312 aga ctc ttt gtt atg gcc agg gaa cat gca cca tcc att ata ttt atg 312 aga ctc ttt gtt atg gcc agg gaa cat gca cca tcc att ata ttt atg 312 aga ctc ttt gtt atg gcc agg atc gcd atca gcc gcd act gcd gcd act act gcd gcd act gcd gcd act act gcd gcd act act gcd act act gcd gcd acc acc acc gcd gcd act gcd gcd acc acc acc gcd gcd acc acc acc acc acc acc acc acc acc a	A:\CBF4\02142003\I462972.raw
299 Leu Wet Dys Turber 120 300 115 300 125 302 ggc ctt gac cag caa att aaa gag atc aaa gag gtc att gag ctt cca 303 Gly Leu Asp Gln Gln Ile Lys Glu Ile Lys Glu Val Ile Glu Leu Pro 303 Gly Leu Asp Gln Gln Ile Lys Glu Ile Lys Glu Val Ile Glu Leu Pro 304 130 306 atc aaa cat ccg gaa ctg ttt gag agc ctt gga att gcg caa cca aag 307 Ile Lys His Pro Glu Leu Phe Glu Ser Leu Gly Ile Ala Gln Pro Lys 307 Ile Lys His Pro Glu Leu Phe Glu Ser Leu Gly Ile Ala Gln Pro Lys 308 145 310 ggt gtc ctt ctt tat gga cct ccg ggc aca gga aag aca ttg ttg gca 310 ggt gtc ctr ctt aca cac act gac tgc acc ttc atc agg gtg tct ggt 311 Gly Val Leu Leu Tyr Gly Pro Pro Gly Thr Gly Lys Thr Leu Leu Ala 311 Gly Val Leu Leu Tyr Gly Pro Pro Gly Thr Phe Ile Arg Val Ser Gly 314 cqt gcg gtt gct cat cac act gac tgc acc ttc atc agg gtg tct ggt 315 Arg Ala Val Ala His His Thr Asp Cys Thr Phe Ile Arg Val Ser Gly 316 180 316 180 316 180 317 180 318 tct gag ttg gtt cag aag tat att ggt gag ggc tcc cgg atg gtt agg 319 Ser Glu Leu Val Gln Lys Tyr Ile Gly Glu Gly Ser Arg Met Val Arg 319 Ser Glu Leu Val Gln Lys Tyr Ile Gly Glu Gly Ser Arg Met Val Arg 319 Ser Glu Leu Phe Val Met Ala Arg Glu His Ala Pro Ser Ile Ile Phe Met 323 Glu Leu Phe Val Met Ala Arg Glu His Ala Pro Ser Ile Ile Phe Met 323 Glu Leu Phe Val Met Ala Arg Glu His Ala Pro Ser Ile Ile Phe Met 324 210 325 240 326 225 330 aac ggt gac agt gaa gtt cag cgt act att gga att gag act ct gaa act gcc 331 Asn Gly Asp Ser Glu Val Gln Arg Thr Met Leu Glu Leu Leu Asn Gln 331 Asn Gly Asp Ser Glu Val Gln Arg Thr Met Leu Glu Leu Leu Asn Gln 332 Leu Asp Gly Phe Glu Ala Ser Asn Lys Ile Lys Val Leu Met Ala Thr 335 Leu Asp Gly Phe Glu Ala Ser Asn Lys Ile Lys Val Leu Met Ala Thr 336 260 376 377 388 2240 389 380 380 380 380 380 380 380 380 380 380	t ata all uuu
303 Gly Leu Rap 130 304 130 306 atc aaa cat cog gaa ctg ttt gag agc ctt gga att gcg caa cca aag 307 Ile Lys His Pro Glu Leu Phe Glu Ser Leu Gly Ile Ala Gln Pro Lys 307 Ile Lys His Pro Glu Leu Phe Glu Ser Leu Gly Ile Ala Gln Pro Lys 308 145 308 145 309 145 310 ggt gtc ctt ctt tat gga cct ccg ggc aca gga aaag aca ttg ttg gca 310 ggt gtc ctt ctt tat gga cct ccg ggc aca gga aaag aca ttg ttg gca 311 Gly Val Leu Leu Tyr Gly Pro Pro Gly Thr Gly Lys Thr Leu Leu Ala 311 Gly Val Leu Leu Tyr Gly Pro Pro Gly Thr Phe Ile Arg Val Ser Gly 314 cgt gcg gtt gct cat cac act gac tgc acc ttc atc agg gtg tct ggt 315 Arg Ala Val Ala His His Thr Asp Cys Thr Phe Ile Arg Val Ser Gly 316 318 tct gag ttg gtc cag aag tat att ggt gag ggc tcc cgg atg gtt agg 317 Ser Glu Leu Val Gln Lys Tyr Ile Gly Glu Gly Ser Arg Met Val Arg 319 Ser Glu Leu Val Gln Lys Tyr Ile Gly Glu Gly Ser Arg Met Val Arg 319 Ser Glu Leu Phe Val Met Ala Arg Glu His Ala Pro Ser Ile Ile Phe Met 323 Glu Leu Phe Val Met Ala Arg Glu His Ala Pro Ser Ile Ile Phe Met 323 Glu Leu Phe Val Met Ala Arg Glu His Ala Pro Ser Ile Ile Phe Met 323 Gaac ggt gac att gac tct atc gga tct gct aga atg gag tct gga act ggc 320 320 220 324 210 326 gat gaa att gac tct atc gga tct gct aga atg gag tct gga act ggc 327 Asp Glu Ile Asp Ser Ile Gly Ser Ala Arg Met Glu Ser Gly Thr Gly 328 225 330 aac ggt gac agt gaa gtt cag cgt act atg ctt gaa ctt cta aac cag 328 225 330 aac ggt gac agt gaa gtt cag cgt act atg ctt gaa ctt cta acc ag 328 226 330 aac ggt gac agt gaa gtt cag cgt act atg ctt gac ctt ctg aga ctt cta acc acc acc acc acc acc acc ac	120 to att gag ctt cca 431
307 The Lys and 145 308	135 caa cca aag 479
311 Giy Val Feu Los 1 165 312 160 313 cgt gcg gtt gct cat cac act gac tgc acc ttc atc agg gtg tct ggt 315 Arg Ala Val Ala His His Thr Asp Cys Thr Phe Ile Arg Val Ser Gly 316 180 318 tct gag ttg gtt cag aag tat att ggt gag ggc tcc cgg atg gtt agg 319 Ser Glu Leu Val Gln Lys Tyr Ile Gly Glu Gly Ser Arg Met Val Arg 319 Ser Glu Leu Val gln Lys Tyr Ile Gly Glu Gly Ser Arg Met Val Arg 319 Ser Glu Leu Val Gln Lys Tyr Ile Gly Glu Gly Ser Arg Met Val Arg 320 320 320 321 322 gaa ctc ttt gtt atg gcc agg gaa cat gca cca tcc att ata ttt atg 323 Glu Leu Phe Val Met Ala Arg Glu His Ala Pro Ser Ile Ile Phe Met 323 Glu Leu Phe Val Met Ala Arg Glu His Ala Pro Ser Ile Ile Phe Met 324 326 gat gaa att gac tct atc gga tct gct aga atg gag tct gga act ggc 327 328 329 320 320 321 320 321 322 323 328 325 330 aac ggt gac agt gaa gtt cag cgt act atg ctt gaa ctt cta aac cag 320 321 322 323 324 325 330 aac ggt gac agt gaa gtt cag cgt act atg ctt gaa ctt cta aac cag 327 328 329 330 aac ggt gac agt gaa gtt cag cgt act atg ctt gaa ctt cta aac cag 331 Asn Gly Asp Ser Glu Val Gln Arg Thr Met Leu Glu Leu Leu Asn Gln 325 330 aac ggt gac agt gaa gca tca aac aaa att aag gtt ttg atg gca acg 331 Asn Gly Asp Ser Glu Val Gln Arg Thr Met Leu Glu Leu Leu Asn Gln 331 Asn Gly Asp Ser Glu Val Gln Arg Thr Met Leu Glu Leu Leu Asn Gln 332 240 334 ctc gat ggt ttt gaa gca tca aac aaa att aag gtt ttg atg gca acg 336 337 338 339 Asn Arg Ile Asp Ile Leu Asp Gln Ala Leu Leu Arg Pro Gly Arg Ile 339 Asn Arg Ile Asp Ile Leu Asp Gln Ala Leu Leu Arg Pro Gly Arg Ile 340 285 340 285 340 285 340 285 340 285 340 285 340 287 340 287 344 345 atc tg aag atc cat tca aga aat atg aac ttg atg cgt ggc att gat 348 349 Asp Arg Lys Ile Glu Phe Pro Asn Pro Asn Glu Asp Ser Arg Phe Asp 340 341 atc ttg aag atc cat tca aga aat atg aac ttg atg cgt ggc att gat 340 340 341 atc tg ag aag atc cat tca aga aat atg aac ttg atg cgt ggc att gat 342 343 344 atc ttg aag atc cat tca aga aat atg aac ttg aga ctc aac acc aac t	150 tag that the gca 527
315 Arg Ala Var 180 316 318 tot gag ttg gtt cag aag tat att ggt gag ggc toc cgg atg gtt agg 318 tot gag ttg gtt cag aag tat att ggt gag ggc toc cgg atg gtt agg 319 Ser Glu Leu Val Gln Lys Tyr Ile Gly Glu Gly Ser Arg Met Val Arg 319 Ser Glu Leu Val Gln Lys Tyr Ile Gly Glu Gly Ser Arg Met Val Arg 320 320 322 gaa ctc ttt gtt atg gcc agg gaa cat gca cca toc att ata ttt atg 323 Glu Leu Phe Val Met Ala Arg Glu His Ala Pro Ser Ile Ile Phe Met 323 Glu Leu Phe Val Met Ala Arg Glu His Ala Pro Ser Ile Ile Phe Met 324 210 326 gat gaa att gac tot atc gga tot gga att gga cot gga act ggc 327 Asp Glu Ile Asp Ser Ile Gly Ser Ala Arg Met Glu Ser Gly Thr Gly 327 Asp Glu Ile Asp Ser Ile Gly Ser Ala Arg Met Glu Ser Gly Thr Gly 328 225 330 aac ggt gac agt gaa gtt cag cgt act atg ott gaa ott ota aac cag 331 Asn Gly Asp Ser Glu Val Gln Arg Thr Met Leu Glu Leu Leu Asn Gln 331 Asn Gly Asp Ser Glu Val Gln Arg Thr Met Leu Glu Leu Leu Asn Gln 331 Asn Gly Asp Ser Glu Val Gln Arg Thr Met Leu Glu Leu Leu Asn Gln 332 240 334 ctc gat ggt tt gaa gca toa aac aaa att aag gtt ttg atg gca acg 335 Leu Asp Gly Phe Glu Ala Ser Asn Lys Ile Lys Val Leu Met Ala Thr 335 Leu Asp Gly Phe Glu Ala Ser Asn Lys Ile Lys Val Leu Met Ala Thr 335 Leu Asp Gly Phe Glu Ala Ser Asn Lys Ile Lys Val Leu Met Ala Thr 336 338 aac aga ata gac att ttg gat caa gcc ott ctg agg cot ggc ata 339 Asn Arg Ile Asp Ile Leu Asp Gln Ala Leu Leu Arg Pro Gly Arg Ile 339 Asn Arg Ile Asp Ile Leu Asp Gln Ala Leu Leu Arg Pro Gly Arg Ile 339 Asn Arg Ile Glu Phe Pro Asn Pro Asn Glu Asp 340 275 342 gac agg aag att gaa ttt coa aat oct aac gag gat toa cgt ttc gat 343 Asp Arg Lys Ile Glu Phe Pro Asn Pro Asn Glu Asp 343 Asp Arg Lys Ile Glu Phe Pro Asn Pro Asn Glu Asp 344 290 346 atc ttg aag atc cat toa aga aaa atg aac ttg atg cgt ggc att gat 347 Ile Leu Lys Ile His Ser Arg Lys Met Asn Leu Met Arg Gly Ile Asp 347 Ile Leu Lys Ile His Ser Arg Lys Met Asn Leu Met Arg Gly Ile Asp	s at a tet agt 5/3
319 Ser Glu lev Val 195 320	185 and att agg 625
323 Glu Leu File 210 324 210 326 gat gaa att gac tct atc gga tct gct aga atg gag tct gga act ggc 327 Asp Glu Ile Asp Ser Ile Gly Ser Ala Arg Met Glu Ser Gly Thr Gly 327 Asp Glu Ile Asp Ser Ile Gly Ser Ala Arg Met Glu Ser Gly Thr Gly 235 328 225 330 aac ggt gac agt gaa gtt cag cgt act atg ctt gaa ctt cta aac cag 331 Asn Gly Asp Ser Glu Val Gln Arg Thr Met Leu Glu Leu Leu Asn Gln 331 Asn Gly Asp Ser Glu Val Gln Arg Thr Met Leu Glu Leu Leu Asn Gln 332 240 334 ctc gat ggt ttt gaa gca tca aac aaa att aag gtt ttg atg gca acg 334 ctc gat ggt ttt gaa gca tca aac aaa att aag gtt ttg atg gca acg 335 Leu Asp Gly Phe Glu Ala Ser Asn Lys Ile Lys Val Leu Met Ala Thr 335 Leu Asp Gly Phe Glu Ala Ser Asn Lys Ile Lys Val Leu Met Ala Thr 336 260 338 aac aga ata gac att ttg gat caa gcc ctt ctg agg cct ggc cgc ata 339 Asn Arg Ile Asp Ile Leu Asp Gln Ala Leu Leu Arg Pro Gly Arg Ile 339 Asn Arg Ile Asp Ile Leu Asp Gln Ala Leu Leu Arg Pro Gly Arg Ile 339 Asp Arg Lys Ile Glu Phe Pro Asn Pro Asn Glu Asp Ser Arg Phe Asp 340 295 344 290 346 atc ttg aag atc cat tca aga aaa atg aac ttg atg cgt ggc att gat 347 Ile Leu Lys Ile His Ser Arg Lys Met Asn Leu Met Arg Gly Ile Asp 347 Ile Leu Lys Ile His Ser Arg Lys Met Asn Leu Met Arg Gly Ile Asp 340 110	200 . Let at at $9/1$
327 Asp Glu lie hop 230 328 225 330 aac ggt gac agt gaa gtt cag cgt act atg ctt gaa ctt cta aac cag 330 aac ggt gac agt gaa gtt cag cgt act atg ctt gaa ctt cta aac cag 331 Asn Gly Asp Ser Glu Val Gln Arg Thr Met Leu Glu Leu Leu Asn Gln 255 250 250 250 250 250 250 250 250 250	215 tet gga act ggc /19
331 Asn Gly Asp Set 245 332 240 334 ctc gat ggt ttt gaa gca tca aac aaa att aag gtt ttg atg gca acg 334 ctc gat ggt ttt gaa gca tca aac aaa att aag gtt ttg atg gca acg 335 Leu Asp Gly Phe Glu Ala Ser Asn Lys Ile Lys Val Leu Met Ala Thr 270 265 336 338 aac aga ata gac att ttg gat caa gcc ctt ctg agg cct ggc cgc ata 339 Asn Arg Ile Asp Ile Leu Asp Gln Ala Leu Leu Arg Pro Gly Arg Ile 285 340 275 340 275 342 gac agg aag att gaa ttt cca aat cct aac gag gat tca cgt ttc gat 342 gac agg aag att gaa ttt cca aat cct aac gag gat tca cgt ttc gat 343 Asp Arg Lys Ile Glu Phe Pro Asn Pro Asn Glu Asp Ser Arg Phe Asp 300 344 290 346 atc ttg aag atc cat tca aga aaa atg aac ttg atg cgt ggc att gat 347 Ile Leu Lys Ile His Ser Arg Lys Met Asn Leu Met Arg Gly Ile Asp 310	230 25 and cag 767
335 Leu Asp Gry The 260 336	230 the ator oca acq 813
339 Asn Arg IIe Nov 280 340 275 342 gac agg aag att gaa ttt cca aat cct aac gag gat tca cgt ttc gat 343 Asp Arg Lys Ile Glu Phe Pro Asn Pro Asn Glu Asp Ser Arg Phe Asp 343 Asp Arg Lys Ile Glu Phe Pro Asn Pro Asn Glu Asp Ser Arg Phe Asp 344 290 344 290 346 atc ttg aag atc cat tca aga aaa atg aac ttg atg cgt ggc att gat 346 atc ttg aag atc cat tca aga aaa atg aac ttg Arg Gly Ile Asp 347 Ile Leu Lys Ile His Ser Arg Lys Met Asn Leu Met Arg Gly Ile Asp 310	265 ant aga aga ata 865
343 Asp Arg Lys 120 295 344 290 346 atc ttg aag atc cat tca aga aaa atg aac ttg atg cgt ggc att gat 346 atc ttg aag atc cat tca aga aaa atg aac ttg atg cgt ggc att gat 347 Ile Leu Lys Ile His Ser Arg Lys Met Asn Leu Met Arg Gly Ile Asp 347 Ile Leu Lys Ile His Ser Arg Lys Met Asn Leu Rga gct gag ctc 1	280 ggt ttc gat
347 He Leu Lys 123 310 and gag act gag ctc 1	295 art aga att gat
	310 aga gct gag ctc 1007
351 Leu Lys Lys Ile Ala Glu Lys Hoo 330 325 352 320 354 aag gcc gtc tgc aca gag gct gga atg ttt gct ctt cgt gag aga agg 1 354 by Ala Val Cys Thr Glu Ala Gly Met Phe Ala Leu Arg Glu Arg Arg 350 345 340 340 350 341 345	325
355 Lys Ala Val Cys Thr Glu Ala Gly 345 345 356 358 gtg cac gtt acc cag gag gac ttc gag atg gca gtg gcc aag gtg atg 358 yval His Val Thr Gln Glu Asp Phe Glu Met Ala Val Ala Lys Val Met 365 360 365	345 La acc aag gtg atg
359 Val His Val Thr Gln Glu Asp Fle 360 360 355 362 aag aaa gac acg gag aag aac atg too otg ogo aag oto tgg aag 362 aag aaa gac acg gag aag aac atg too otg ogo aag oto tgg	g aag aac atg tcc ctg cgc aag ctc tgg aag 1148

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/09/462,972 DATE: 02/14/2003 TIME: 11:52:51

Input Set : A:\BB-1095-A Corrected Seq List.txt

Output Set: N:\CRF4\02142003\I462972.raw

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:11; N Pos. 262,271,293,331,355,356,373,384,387,391,398,405,407,418,425

Seq#:11; N Pos. 427,428,453,454,473,497,518,526,527,541

Seq#:13; N Pos. 149,206,217,299,325,354,412,478,493,498,521,558,571,584,585

Seq#:14; Xaa Pos. 20,47,66,70,97,101,106,115,116

DATE: 02/14/2003 VERIFICATION SUMMARY TIME: 11:52:51 PATENT APPLICATION: US/09/462,972

Input Set : A:\BB-1095-A Corrected Seq List.txt

Output Set: N:\CRF4\02142003\I462972.raw

L:34 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:32 L:269 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:3,Line#:267 L:1222 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:240

M:341 Repeated in SeqNo=11

L:1283 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:64

M:341 Repeated in SeqNo=12

L:1393 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:120

L:1469 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:16

M:341 Repeated in SeqNo=14